



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

10/8/6, ST1
JFW
4/8/84

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10816591

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



IFWO

RAW SEQUENCE LISTING

DATE: 04/08/2004

PATENT APPLICATION: US/10/816,591

TIME: 10:13:52

Input Set : A:\NHL-NP-46 Sequence Listing.txt

Output Set: N:\CRF4\04082004\J816591.raw

3 <110> APPLICANT: Fuertes-Lopez, Laura
 4 Timon-Jimenez, Marcos
 6 <120> TITLE OF INVENTION: DNA EXPRESSION CONSTRUCT FOR TREATMENT OF INFECTIONS WITH
 7 LEISHMANIASIS
 9 <130> FILE REFERENCE: NHL-NP-46
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/816,591
 C--> 11 <141> CURRENT FILING DATE: 2004-04-01
 11 <150> PRIOR APPLICATION NUMBER: DE 101 48 732.0
 12 <151> PRIOR FILING DATE: 2001-10-02
 14 <150> PRIOR APPLICATION NUMBER: DE 101 56 679.4
 15 <151> PRIOR FILING DATE: 2001-11-12
 17 <150> PRIOR APPLICATION NUMBER: PCT/DE02/03799
 18 <151> PRIOR FILING DATE: 2002-10-02
 20 <160> NUMBER OF SEQ ID NOS: 9
 22 <170> SOFTWARE: PatentIn version 3.2
 24 <210> SEQ ID NO: 1
 25 <211> LENGTH: 4791
 26 <212> TYPE: DNA
 27 <213> ORGANISM: Plasmid pSCp36
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 31 <221> NAME/KEY: misc_feature
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 33 <223> OTHER INFORMATION: Kanamycin resistance, reverse complementary
 35 <220> FEATURE:
 36 <221> NAME/KEY: promoter
 37 <222> LOCATION: (2447)..(3243)
 38 <223> OTHER INFORMATION: CMV
 40 <220> FEATURE:
 41 <221> NAME/KEY: Intron
 42 <222> LOCATION: (3250)..(3386)
 44 <220> FEATURE:
 45 <221> NAME/KEY: misc_feature
 46 <222> LOCATION: (3393)..(4331)
 47 <223> OTHER INFORMATION: p36 protein
 49 <220> FEATURE:
 50 <221> NAME/KEY: polyA_site
 51 <222> LOCATION: (4338)..(4539)
 52 <223> OTHER INFORMATION: poly A site aus SV 40
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 57 tcagctcact caaaggcggg aatacggtta tccacagaat caggggataa cgcaggaaag 120
 59 aacatgtctc gggaggctc acgtgacatg tgagcaaaa gccagcaaaa ggccaggaaac 180
 61 cgtaaaaagg ccgcgttgct ggcgtttttc cataggctcc gccccctga cgagcatcac 240

Does Not Comply
 Corrected Diskette Needed
 CP5.3-4

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Input Set : A:\NHL-NP-46 Sequence Listing.txt

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69	ctcagttcgg	tgtaggtcgt	tcgctccaag	ctgggctgtg	tgcacgaacc	ccccgttcag	480
71	cccgaccgct	gcgccttatc	cgtaactat	cgtcttgagt	ccaacccggt	aagacacgac	540
73	ttatcgccac	tggcagcagc	cactggtaac	aggattagca	gagcgaggta	tgtaggcggg	600
75	gctacagagt	tcttgaagtg	gtggcctaac	tacggctaca	ctagaaggac	agtatttggg	660
77	atctgcgctc	tgctgaagcc	agttaccttc	ggaaaaagag	ttggtagctc	ttgatccggc	720
79	aaacaaacca	ccgctggtag	cggtggtttt	tttgtttgca	agcagcagat	tacgcgcaga	780
81	aaaaaaggat	ctcaagaaga	tcctttgatc	ttttctacgg	ggtctgacgc	tcagtggaac	840
83	gaaaactcac	gttaagggat	tttggtcagt	agattatcaa	aaaggatctt	cacctagatc	900
85	cttttaaat	aaaaatgaag	ttttaaatca	atctaaagta	tatatgagta	aacttggctc	960
87	gacagttaac	aatgcttaac	cagtgaagca	cctatctcag	cgatctgtct	atttcgttca	1020
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91	gcgctgcgaa	tcgggagcgg	cgataccgta	aagcacgagg	aagcggtcag	cccattcgcc	1140
93	gccaagctct	tcagcaatat	cacgggtagc	caacgctatg	tcctgatagc	ggtccgccac	1200
95	accagccgg	ccacagtcga	tgaatccaga	aaagcggcca	ttttccacca	tgatattcgg	1260
97	caagcaggca	tcgccatggg	tcacgacgag	atcctcgccg	tcgggcatgc	tcgccttgag	1320
99	cctggcgaac	agttcggctg	gcgcgagccc	ctgatgctct	tcgtccagat	catcctgatc	1380
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117	agggttattg	tctcatgagc	ggatacatat	ttgaatgtat	ttagaaaaat	aaacaaatag	1920
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131	tcgcgaactg	ttgggaaggg	cgatcgggtg	gggcctcttc	gctattacgc	cagctggcga	2340
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143	gcccataat	ggagtccgc	gttacataac	ttacggtaaa	tggcccgcct	ggctgaccgc	2700
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147	ggactttcca	ttgacgtcaa	tgggtggagt	atttacggta	aactgccac	ttggcagtac	2820
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151	cctggcatta	tgcccagtac	atgaccttac	gggactttcc	tacttggcag	tacatctacg	2940
153	tattagtcat	cgctattacc	atggtgatgc	ggttttggca	gtacaccaat	gggcgtggat	3000
155	agcggtttga	ctcacgggga	tttccaagtc	tcacccccat	tgacgtcaat	gggagtttgt	3060
157	tttggcacca	aaatcaacgg	gactttccaa	aatgtcgtaa	taaccccgcc	ccgttgacgc	3120
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TIME: 10:13:52

Input Set : A:\NHL-NP-46 Sequence Listing.txt

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```

161 gtcagatcac tagaagcttt attgcggttag tttatcacag ttaaattgct aacgcagtca 3240
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165 ggcttgctga gacagagaag actcttgctg ttctgatagg cacctattgg tcttactgac 3360
167 atccactttg cctttctctc cacaggggta ccatgaacta cgagggtcac ctgaagggcc 3420
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191 tcaagatcaa cgtggagtgc cccatcaacc agatgcctt ctgcaccaac cgcttctgga 4140
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203 aacaagttaa caacaacaat tgcattcatt ttatgtttca ggttcagggg gaggtgtggg 4500
205 aggtttttta aagcaagtaa aacctctaca aatgtggtag aattcagggg gagacccaat 4560
207 tcgtaatcat ggtcatagct gtttcctgtg tgaaattgtt atccgctcac aattccacac 4620
209 aacatacgag ccggaagcat aaagtgtaaa gcctgggggtg cctaattgagt gagctaactc 4680
211 acattaattg cgttcgctc actgcccgct ttccagtcgg gaaacctgtc gtgccagctg 4740
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```

216 <210> SEQ ID NO: 2

217 <211> LENGTH: 11

218 <212> TYPE: PRT

219 <213> ORGANISM: Human immunodeficiency virus type 1, TAT peptide

221 <400> SEQUENCE: 2

223 Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg

224 1 5 10

227 <210> SEQ ID NO: 3

228 <211> LENGTH: 7

229 <212> TYPE: PRT

230 <213> ORGANISM: Simian virus 40, NLS peptide

232 <400> SEQUENCE: 3

234 Pro Lys Lys Lys Arg Lys Val

235 1 5

238 <210> SEQ ID NO: 4

239 <211> LENGTH: 20

240 <212> TYPE: DNA

241 <213> ORGANISM: ODN 1

244 <220> FEATURE:

245 <221> NAME/KEY: modified base

246 <222> LOCATION: (12)..(12)

247 <223> OTHER INFORMATION: xT = Thymin modified with a reactive amino group

mandatory, <213>
 Response has to be
 either Artificial/
 Unknown or
 Genus/
 Species.

Please see item
 #10 on error Summary
 Sheet.

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/816,591

DATE: 04/08/2004

TIME: 10:13:52

Input Set : A:\NHL-NP-46 Sequence Listing.txt

Output Set: N:\CRF4\04082004\J816591.raw

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250 gggagtccag ttttctggac
253 <210> SEQ ID NO: 5
254 <211> LENGTH: 20
255 <212> TYPE: DNA
256 <213> ORGANISM: ODN 2
258 <400> SEQUENCE: 5
259 aggggtccag ttttctggac
262 <210> SEQ ID NO: 6
263 <211> LENGTH: 33
264 <212> TYPE: DNA
265 <213> ORGANISM: 1. PCR: Primer left
267 <400> SEQUENCE: 6
268 ttatatggtta ccatgaacat acgaggggtca cct
271 <210> SEQ ID NO: 7
272 <211> LENGTH: 42
273 <212> TYPE: DNA
274 <213> ORGANISM: 1. PCR: Primer right
276 <400> SEQUENCE: 7
277 ttatatgagc tcagaagaca cggacaggga cctcttccgt cg
280 <210> SEQ ID NO: 8
281 <211> LENGTH: 33
282 <212> TYPE: DNA
283 <213> ORGANISM: 2. PCR: Primer left
285 <400> SEQUENCE: 8
286 ttatatggtta ccatgaacat acgaggggtca cct
289 <210> SEQ ID NO: 9
290 <211> LENGTH: 33
291 <212> TYPE: DNA
292 <213> ORGANISM: 2. PCR: Primer right
294 <400> SEQUENCE: 9
295 ttatatgagc tcttactcgg ccgtcggaga tgg

Some errors
See item #10 on erran Summary Sheet

VERIFICATION SUMMARY

DATE: 04/08/2004

PATENT APPLICATION: US/10/816,591

TIME: 10:13:53

Input Set : A:\NHL-NP-46 Sequence Listing.txt

Output Set: N:\CRF4\04082004\J816591.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date